RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	
Source:	1FW/6
Date Processed by STIC:	12/30/05
·	

ENTERED



IFW16

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/955,502A

DATE: 12/30/2005
TIME: 09:32:37

Input Set : A:\960296.97559.txt

Output Set: N:\CRF4\12302005\I955502A.raw

3 <110> APPLICANT: Downs, Diana M. Gralnick, Jeff A. 6 <120> TITLE OF INVENTION: Method for Preventing Superoxide Damage to Cells and Oxygen-Labile Proteins 9 <130> FILE REFERENCE: 960296.97559 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/955,502A C--> 11 <141> CURRENT FILING DATE: 2001-09-18 11 <160> NUMBER OF SEQ ID NOS: 45 13 <170> SOFTWARE: PatentIn version 3.3 15 <210> SEQ ID NO: 1 16 <211> LENGTH: 65 17 <212> TYPE: PRT 18 <213> ORGANISM: Artificial 20 <220> FEATURE: 21 <223> OTHER INFORMATION: synthetic YggX consensus sequence 24 <220> FEATURE: 25 <221> NAME/KEY: UNSURE 26 <222> LOCATION: (2)..(2) 27 <223> OTHER INFORMATION: can be any amino acid 29 <220> FEATURE: 30 <221> NAME/KEY: UNSURE 31 <222> LOCATION: (4)..(6) 32 <223> OTHER INFORMATION: can be any amino acid 34 <220> FEATURE: 35 <221> NAME/KEY: UNSURE 36 <222> LOCATION: (8)..(22) 37 <223> OTHER INFORMATION: can be any amino acid 39 <220> FEATURE: 40 <221> NAME/KEY: UNSURE 41 <222> LOCATION: (24)..(26) 42 <223> OTHER INFORMATION: can be any amino acid 44 <220> FEATURE: 45 <221> NAME/KEY: UNSURE 46 <222> LOCATION: (28)..(38) 47 <223> OTHER INFORMATION: can be any amino acid 49 <220> FEATURE: 50 <221> NAME/KEY: UNSURE 51 <222> LOCATION: (40)..(41) 52 <223> OTHER INFORMATION: can be any amino acid 54 <220> FEATURE: 55 <221> NAME/KEY: UNSURE 56 <222> LOCATION: (43)..(45)

57 <223> OTHER INFORMATION: can be any amino acid

Input Set : A:\960296.97559.txt

Output Set: N:\CRF4\12302005\I955502A.raw

```
59 <220> FEATURE:
     60 <221> NAME/KEY: UNSURE
     61 <222> LOCATION: (48)..(48)
     62 <223> OTHER INFORMATION: can be any amino acid
     64 <220> FEATURE:
     65 <221> NAME/KEY: UNSURE
     66 <222> LOCATION: (50)..(50)
     67 <223> OTHER INFORMATION: can be any amino acid
     69 <220> FEATURE:
     70 <221> NAME/KEY: UNSURE
     71 <222> LOCATION: (53)..(54)
     72 <223> OTHER INFORMATION: can be any amino acid
     74 <220> FEATURE:
     75 <221> NAME/KEY: UNSURE
     76 <222> LOCATION: (56)..(62)
     77 <223> OTHER INFORMATION: can be any amino acid
     79 <220> FEATURE:
     80 <221> NAME/KEY: UNSURE
     81 <222> LOCATION: (64)..(65)
     82 <223> OTHER INFORMATION: can be any amino acid
     84 <400> SEQUENCE: 1
W--> 86 Met Xaa Arg Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                        5
     87 1
                                             10
                                                                 15
W--> 90 Xaa Xaa Xaa Xaa Xaa Xaa Pro Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa
                    20
                                         25
W--> 94 Xaa Xaa Xaa Xaa Xaa Xaa Trp Xaa Xaa Trp Xaa Xaa Xaa Gln Thr Xaa
                                    40
     95
                35
W--> 98 Leu Xaa Asn Glu Xaa Xaa Leu Xaa Xaa Xaa Xaa Xaa Xaa Arg Xaa
     99
            50
W--> 102 Xaa
     103 65
     106 <210> SEQ ID NO: 2
     107 <211> LENGTH: 87
     108 <212> TYPE: PRT
     109 <213> ORGANISM: Bordetella pertussis
     111 <400> SEQUENCE: 2
     113 Met Ser Arg Ile Val Asn Cys Val Lys Leu Lys Arg Glu Ala Glu Gly
     114 1
     117 Leu Asp Phe Pro Pro Tyr Pro Gly Glu Leu Gly Thr Arg Ile Trp Gln
     118
                     20
                                          25
     121 Gln Ile Ser Lys Glu Ala Trp Glu Glu Trp Lys Gln Ile Gln Thr Arg
     122
     125 Leu Val Asn Glu Asn Arg Leu Asn Leu Ala Asp Ala Arg Ala Arg Lys
     126
             50
                                 55
     129 Tyr Leu Gln Gln Met Glu Arg Phe Leu Phe Glu Asp Gly Thr Val
     130 65
                             70
                                                  75
                                                                      80
     133 Glu Ala Gln Gly Tyr Val Pro
     134
     137 <210> SEQ ID NO: 3
```

Input Set : A:\960296.97559.txt

Output Set: N:\CRF4\12302005\I955502A.raw

138 <211> LENGTH: 87 139 <212> TYPE: PRT 140 <213> ORGANISM: Bordetella parapertussis 142 <400> SEQUENCE: 3 144 Met Ser Arg Ile Val Asn Cys Val Lys Leu Lys Arg Glu Ala Glu Gly 145 1 148 Leu Asp Phe Pro Pro Tyr Pro Gly Glu Leu Gly Thr Arg Ile Trp Gln 149 20 30 152 Gln Ile Ser Lys Glu Ala Trp Glu Glu Trp Lys Gln Ile Gln Thr Arg 153 35 40 45 156 Leu Val Asn Glu Asn Arg Leu Asn Leu Ala Asp Ala Arg Ala Arg Lys 157 55 160 Tyr Leu Gln Gln Met Glu Arg Phe Leu Phe Glu Asp Gly Thr Val 161 65 70 75 80 164 Glu Ala Gln Gly Tyr Val Pro 165 168 <210> SEQ ID NO: 4 169 <211> LENGTH: 86 170 <212> TYPE: PRT 171 <213> ORGANISM: Bordetella bronchiseptica 173 <400> SEQUENCE: 4 175 Met Ser Arg Ile Val Asn Cys Val Lys Leu Lys Arg Glu Ala Glu Gly 176 1 5 10 15 179 Leu Asp Phe Pro Pro Tyr Pro Gly Glu Leu Gly Thr Arg Ile Trp Gln 180 183 Gln Ile Ser Lys Glu Ala Trp Glu Glu Trp Lys Gln Ile Gln Thr Arq 184 35 40 187 Leu Val Asn Glu Asn Arg Leu Asn Leu Ala Asp Ala Arg Ala Arg Lys 188 50 55 191 Tyr Leu Gln Gln Met Glu Arg Phe Leu Phe Glu Asp Gly Thr Val 192 65 70 75 80 195 Glu Ala Gln Gly Val Pro 199 <210> SEQ ID NO: 5 200 <211> LENGTH: 91 201 <212> TYPE: PRT 202 <213> ORGANISM: Actinobacillus actinomycetemcomitans 204 <400> SEQUENCE: 5 206 Met Ala Arg Met Val Phe Cys Glu Arg Leu Lys Gln Glu Ala Glu Gly 207 1 10 210 Leu Asp Phe Gln Leu Tyr Pro Gly Glu Leu Gly Lys Arg Ile Phe Asp 211 20 214 Ser Ile Ser Lys Gln Ala Trp Gly Glu Trp Met Lys Lys Gln Thr Met 215 35 218 Leu Val Asn Glu Lys Lys Leu Asn Met Met Asn Ala Glu His Arg Lys 219 50 55 60 222 Leu Leu Glu Gln Glu Met Val Asn Phe Leu Phe Glu Gly Lys Asp Val 70 226 His Ile Glu Gly Tyr Thr Pro Pro Glu Ala Lys

Input Set : A:\960296.97559.txt

Output Set: N:\CRF4\12302005\I955502A.raw

```
227
                                         90
                    85
230 <210> SEQ ID NO: 6
231 <211> LENGTH: 87
232 <212> TYPE: PRT
233 <213> ORGANISM: Pasteurella multocida
235 <400> SEQUENCE: 6
237 Met Ala Arg Thr Val Phe Cys Glu Tyr Leu Lys Gln Glu Ser Glu Gly
238 1
                    5
                                         10
                                                              15
241 Leu Asp Phe Gln Leu Tyr Pro Gly Glu Leu Gly Lys Arg Ile Phe Asp
242
                20
                                     25
                                                          30
245 Ser Ile Ser Lys Gln Ala Trp Arg Glu Trp Met Lys Lys Gln Thr Met
249 Leu Val Asn Glu Lys Lys Leu Asn Met Met Asn Ala Asp His Arg Gln
250
        50
                             55
253 Leu Leu Glu Gln Glu Met Val Asn Phe Leu Phe Glu Gly Lys Asp Val
254 65
                        70
                                             75
                                                                  80
257 His Ile Glu Gly Tyr Val Pro
258
                    85
261 <210> SEQ ID NO: 7
262 <211> LENGTH: 87
263 <212> TYPE: PRT
264 <213 > ORGANISM: Haemophilus influenzae
266 <400> SEQUENCE: 7
268 Met Ala Arg Thr Val Phe Cys Glu Tyr Leu Lys Lys Glu Ala Glu Gly
269 1
272 Leu Asp Phe Gln Leu Tyr Pro Gly Glu Leu Gly Lys Arg Ile Phe Asp
273
                20
                                                          30
                                     25
276 Ser Val Ser Lys Gln Ala Trp Gly Glu Trp Ile Lys Lys Gln Thr Met
            35
                                 40
277
280 Leu Val Asn Glu Lys Lys Leu Asn Met Met Asn Ala Glu His Arg Lys
281
        50
                             55
284 Leu Leu Glu Gln Glu Met Val Asn Phe Leu Phe Glu Gly Lys Asp Val
288 His Ile Glu Gly Tyr Val Pro
289
292 <210> SEQ ID NO: 8
293 <211> LENGTH: 87
294 <212> TYPE: PRT
295 <213> ORGANISM: Haemophilus ducreyi
297 <400> SEQUENCE: 8
299 Met Ala Arg Met Val Phe Cys Glu Tyr Leu Lys Lys Glu Ala Glu Gly
300 1
303 Leu Asp Phe Gln Leu Tyr Pro Gly Glu Leu Gly Lys Arg Ile Phe Asn
                20
304
                                     25
                                                          30
307 Ser Ile Ser Lys Gln Ala Trp Ala Glu Trp Ile Lys Lys Gln Thr Met
308
            35
                                 40
311 Leu Val Asn Glu Lys Lys Leu Asn Met Met Asn Pro Glu His Arg Gln
                             55
312
315 Leu Leu Glu Ala Glu Met Val Asn Phe Leu Phe Glu Gly Lys Asp Val
```

Input Set : A:\960296.97559.txt

Output Set: N:\CRF4\12302005\I955502A.raw

```
316 65
                        70
                                             75
                                                                  80
319 His Ile Asp Gly Tyr Val Pro
320
                    85
323 <210> SEQ ID NO: 9
324 <211> LENGTH: 88
325 <212> TYPE: PRT
326 <213> ORGANISM: Shewanella putrefaciens
328 <400> SEQUENCE: 9
330 Met Ala Arg Thr Val Asn Cys Val His Leu Asn Lys Glu Ala Asp Gly
                    5
331 1
                                         10
                                                              15
334 Leu Asp Phe Gln Leu Tyr Pro Gly Asp Leu Gly Lys Arg Ile Phe Asp
                                     25
335
338 Asn Ile Ser Lys Glu Ala Trp Gly Leu Trp Gln Lys Lys Gln Thr Met
339
            35
                                 40
342 Leu Ile Asn Glu Lys Lys Leu Asn Met Met Asn Val Asp Asp Arg Lys
343
        50
                             55
346 Phe Leu Glu Ala Gln Met Thr Ser Phe Leu Phe Glu Gly Lys Asp Val
347 65
                        70
                                             75
                                                                  80
350 Glu Ile Glu Gly Phe Val Pro Glu
351
                    85
354 <210> SEQ ID NO: 10
355 <211> LENGTH: 90
356 <212> TYPE: PRT
357 <213> ORGANISM: Vibrio cholerae
359 <400> SEQUENCE: 10
361 Met Ala Arg Thr Val Phe Cys Thr Arg Leu Gln Lys Glu Ala Asp Gly
                    5
362 1
                                         10
365 Leu Asp Phe Gln Leu Tyr Pro Gly Glu Leu Gly Lys Arg Ile Phe Asp
366
                20
                                     25
369 Asn Ile Cys Lys Glu Ala Trp Ala Gln Trp Gln Thr Lys Gln Thr Met
            35
370
373 Leu Ile Asn Glu Lys Lys Leu Asn Met Met Asp Pro Glu His Arg Lys
                        55
377 Leu Leu Glu Gln Glu Met Val Asn Phe Leu Phe Glu Gly Lys Glu Val
                        70
                                             75
381 His Ile Glu Gly Tyr Thr Pro Pro Ala Lys
382
                    85
                                         90
385 <210> SEQ ID NO: 11
386 <211> LENGTH: 91
387 <212> TYPE: PRT
388 <213> ORGANISM: Escherichia coli K-12 MG1655
390 <400> SEQUENCE: 11
392 Met Ser Arg Thr Ile Phe Cys Thr Phe Leu Gln Arg Glu Ala Glu Gly
393 1
                                                              15
396 Gln Asp Phe Gln Leu Tyr Pro Gly Glu Leu Gly Lys Arg Ile Tyr Asn
397
                20
                                     25
                                                          30
400 Glu Ile Ser Lys Glu Ala Trp Ala Gln Trp Gln His Lys Gln Thr Met
404 Leu Ile Asn Glu Lys Lys Leu Asn Met Met Asn Ala Glu His Arg Lys
```

Input Set : A:\960296.97559.txt

Output Set: N:\CRF4\12302005\I955502A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 2,4,5,6,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,24,25 Seq#:1; Xaa Pos. 26,28,29,30,31,32,33,34,35,36,37,38,40,41,43,44,45,48,50 Seq#:1; Xaa Pos. 53,54,56,57,58,59,60,61,62,64,65

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1

VERIFICATION SUMMARY DATE: 12/30/2005

PATENT APPLICATION: US/09/955,502A TIME: 09:32:38

Input Set : A:\960296.97559.txt

Output Set: N:\CRF4\12302005\1955502A.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:86 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:16

L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:32

L:98 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:48

L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:64